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ed.
1 to 2384)
Lishko,V.R.
S 561508-A-1 24-JUN-1997;
Location/Qualifiers
..2384
Organism="unknown"
533 c 520 g 679 t
100.0%; Score 300; DB 6; Length 2384;
Key 100.0%; Pred.No.3.2e-58;
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osinase (TYR) mRNA, complete cds.
2384 bp mRNA linear PRI 0
GI:1698397
ens
Metazoa; Chordata; Craniata; Vertebrata; Eutele
Euthelia; Primates; Catarrhini; Homnidae; Homo
..S., Tomita.Y., Tagami.H., Muller,R.M. and Cohen
basis for the heterogeneity of human tyrosinase
Exp. Med. 156 (4), 403-414 (1988)
1 to 600)
Tomita,Y., Okinaga,S., Tagami,H. and Shibahara
analysis of the cDNA encoding human tyrosinase
Biophys. Res. Commun. 162 (3), 984-990 (1989)
1996 this sequence version replaced gi:340036.
arry and computer-readable copy of sequence [1] Kikkawa
by S.Shibahara, 15-AUG-1989.
Location/Qualifiers
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cell_line="Memo melanoma DNA, and cDNA to mRNA
tissue_type="Placenta"
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Matches 300; Conservative 0; Mismatches 0; Indels 0; Gaps 0		
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DB 244	GGTGAATAGAGTTTTCACCTTAACATTAAGCCCTATCCACCTGTGGATACGACCAAT 303	
QY 181	cgaagaagaatcagtcacatgctcttccacagagatgaaagcttaagataaaagactaaag 240	
DB 304	CGAAGAAAAAGTCAGTCAGTCTTTTCAGAGAGATGAAGAGCTTAAGATMAAGACTTAAAG 363	
QY 241	tcatttgcctggaaggttgagtggtattatataagctcgaagcaagcatgtgataatc 300	
DB 364	TGATTTCATGCTGAGGTGGAGTGTATTATTATAGGCTCTCAGCCACATGTGTATATC 423	
RESULT 6		
DBD 124	TCATTTCGCAAGGTCAAAATCATTATGTTTGTAGTCTATTAACTGGGTTGCTTAAGTC 183	
DB 184	AGCCATTATTATTACTAACTTATGTATATTTCTAAACCTAAGCAATTAACATTAAT 243	
DB 121	ggtgaatagagtttctcattacaataggccatcccaactgtgtggatagagccaatt 180	
DB 244	GGTGAATAGAGTTTTCACCTTAACATTAAGCCCTATCCACCTGTGGATACGACCAAT 303	
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DB 121	ggtgaatagagtttctcattacaataggccatcccaactgtgtggatagagccaatt 180	
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RESULT 6		
DBD 124	TCATTTCGCAAGGTCAAAATCATTATGTTTGTAGTCTATTAACTGGGTTGCTTAAGTC 183	
DB 184	AGCCATTATTATTACTAACTTATGTATATTTCTAAACCTAAGCAATTAACATTAAT 243	
DB 121	ggtgaatagagtttctcattacaataggccatcc	